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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/887,669

DATE: 04/15/2003

TIME: 13:17:22

Input Set : A:\38621246.app

Output Set: N:\CRF4\04152003\I887669.raw

3 <110> APPLICANT: SCHLESSINGER, JOSEPH  
4 SAP, JAN M.  
5 ULLRICH, AXEL  
6 VOGEL, WOLFGANG  
7 FUCHS, MIRIAM  
9 <120> TITLE OF INVENTION: NOVEL RECEPTOR-TYPE PHOSPHOTYROSINE PHOSPHATASE-KAPPA  
11 <130> FILE REFERENCE: 038602/1246  
13 <140> CURRENT APPLICATION NUMBER: 09/887,669  
C--> 14 <141> CURRENT FILING DATE: 2001-10-10  
16 <150> PRIOR APPLICATION NUMBER: 09/234,883  
17 <151> PRIOR FILING DATE: 1999-01-21  
19 <150> PRIOR APPLICATION NUMBER: 08/087,244  
20 <151> PRIOR FILING DATE: 1993-07-01  
22 <150> PRIOR APPLICATION NUMBER: 08/049,384  
23 <151> PRIOR FILING DATE: 1993-04-21  
25 <160> NUMBER OF SEQ ID NOS: 13  
27 <170> SOFTWARE: PatentIn Ver. 2.1  
29 <210> SEQ ID NO: 1  
30 <211> LENGTH: 1457  
31 <212> TYPE: PRT  
32 <213> ORGANISM: Mus musculus  
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39 20 25 30  
41 Gly Gly Cys Thr Phe Asp Asp Gly Pro Gly Ala Cys Asp Tyr His Gln  
42 35 40 45  
44 Asp Leu Tyr Asp Asp Phe Glu Trp Val His Val Ser Ala Gln Glu Pro  
45 50 55 60  
47 His Tyr Leu Pro Pro Glu Met Pro Gln Gly Ser Tyr Met Val Val Asp  
48 65 70 75 80  
50 Ser Ser Asn His Asp Pro Gly Glu Lys Ala Arg Leu Gln Leu Pro Thr  
51 85 90 95  
53 Met Lys Glu Asn Asp Thr His Cys Ile Asp Phe Ser Tyr Leu Leu Tyr  
54 100 105 110  
56 Ser Gln Lys Gly Leu Asn Pro Gly Thr Leu Asn Ile Leu Val Arg Val  
57 115 120 125  
59 Asn Lys Gly Pro Leu Ala Asn Pro Ile Trp Asn Val Thr Gly Phe Thr  
60 130 135 140  
62 Gly Arg Asp Trp Leu Arg Ala Glu Leu Ala Val Ser Thr Phe Trp Pro  
63 145 150 155 160  
65 Asn Glu Tyr Gln Val Ile Phe Glu Ala Glu Val Ser Gly Gly Arg Ser

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66          165          170          175
68 Gly Tyr Ile Ala Ile Asp Asp Ile Gln Val Leu Ser Tyr Pro Cys Asp
69          180          185          190
71 Lys Ser Pro His Phe Leu Arg Leu Gly Asp Val Glu Val Asn Ala Gly
72          195          200          205
74 Gln Asn Ala Thr Phe Gln Cys Ile Ala Thr Gly Arg Asp Ala Val His
75          210          215          220
77 Asn Lys Leu Trp Leu Gln Arg Arg Asn Gly Glu Asp Ile Pro Val Ala
78 225          230          235          240
80 Gln Thr Lys Asn Ile Asn His Arg Arg Phe Ala Ala Ser Phe Arg Leu
81          245          250          255
83 Gln Glu Val Thr Lys Thr Asp Gln Asp Leu Tyr Arg Cys Val Thr Gln
84          260          265          270
86 Ser Glu Arg Gly Ser Gly Val Ser Asn Phe Ala Gln Leu Ile Val Arg
87          275          280          285
89 Glu Pro Pro Arg Pro Ile Ala Pro Pro Gln Leu Leu Gly Val Gly Pro
90          290          295          300
92 Thr Tyr Leu Leu Ile Gln Leu Asn Ala Asn Ser Ile Ile Gly Asp Gly
93 305          310          315          320
95 Pro Ile Ile Leu Lys Glu Val Glu Tyr Arg Met Thr Ser Gly Ser Trp
96          325          330          335
98 Thr Glu Thr His Ala Val Asn Ala Pro Thr Tyr Lys Leu Trp His Leu
99          340          345          350
101 Asp Pro Asp Thr Glu Tyr Glu Ile Arg Val Leu Leu Thr Arg Pro Gly
102          355          360          365
104 Glu Gly Gly Thr Gly Leu Pro Gly Pro Pro Leu Ile Thr Arg Thr Lys
105          370          375          380
107 Cys Ala Glu Pro Met Arg Thr Pro Lys Thr Leu Lys Ile Ala Glu Ile
108 385          390          395          400
110 Gln Ala Arg Arg Ile Ala Val Asp Trp Glu Ser Leu Gly Tyr Asn Ile
111          405          410          415
113 Thr Arg Cys His Thr Phe Asn Val Thr Ile Cys Tyr His Tyr Phe Arg
114          420          425          430
116 Gly His Asn Glu Ser Arg Ala Asp Cys Leu Asp Met Asp Pro Lys Ala
117          435          440          445
119 Pro Gln His Val Val Asn His Leu Pro Pro Tyr Thr Asn Val Ser Leu
120          450          455          460
122 Lys Met Ile Leu Thr Asn Pro Glu Gly Arg Lys Glu Ser Glu Glu Thr
123 465          470          475          480
125 Ile Ile Gln Thr Asp Glu Asp Val Pro Gly Pro Val Pro Val Lys Ser
126          485          490          495
128 Leu Gln Gly Thr Ser Phe Glu Asn Lys Ile Phe Leu Asn Trp Lys Glu
129          500          505          510
131 Pro Leu Glu Pro Asn Gly Ile Ile Thr Gln Tyr Glu Val Ser Tyr Ser
132          515          520          525
134 Ser Ile Arg Ser Phe Asp Pro Ala Val Pro Val Ala Gly Pro Pro Gln
135          530          535          540
137 Thr Val Ser Asn Leu Trp Asn Ser Thr His His Val Phe Met His Leu
138 545          550          555          560

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140 His Pro Gly Thr Thr Tyr Gln Phe Phe Ile Arg Ala Ser Thr Val Lys
141                               565                               570                               575
143 Gly Phe Gly Pro Ala Thr Ala Ile Asn Val Thr Thr Asn Ile Ser Ala
144                               580                               585                               590
146 Pro Ser Leu Pro Asp Tyr Glu Gly Val Asp Ala Ser Leu Asn Glu Thr
147                               595                               600                               605
149 Ala Thr Thr Ile Thr Val Leu Leu Arg Pro Ala Gln Ala Lys Gly Ala
150                               610                               615                               620
152 Pro Ile Ser Ala Tyr Gln Ile Val Val Glu Gln Leu His Pro His Arg
153 625                               630                               635                               640
155 Thr Lys Arg Glu Ala Gly Ala Met Glu Cys Tyr Gln Val Pro Val Thr
156                               645                               650                               655
158 Tyr Gln Asn Ala Leu Ser Gly Gly Ala Pro Tyr Tyr Phe Ala Ala Glu
159                               660                               665                               670
161 Leu Pro Pro Gly Asn Leu Pro Glu Pro Ala Pro Phe Thr Val Gly Asp
162                               675                               680                               685
164 Asn Arg Thr Tyr Lys Gly Phe Trp Asn Pro Pro Leu Ala Pro Arg Lys
165                               690                               695                               700
167 Gly Tyr Asn Ile Tyr Phe Gln Ala Met Ser Ser Val Glu Lys Glu Thr
168 705                               710                               715                               720
170 Lys Thr Gln Cys Val Arg Ile Ala Thr Lys Ala Ala Ala Thr Glu Glu
171                               725                               730                               735
173 Pro Glu Val Ile Pro Asp Pro Ala Lys Gln Thr Asp Arg Val Val Lys
174                               740                               745                               750
176 Ile Ala Gly Ile Ser Ala Gly Ile Leu Val Phe Ile Leu Leu Leu Leu
177                               755                               760                               765
179 Val Val Ile Val Ile Val Lys Lys Ser Lys Leu Ala Lys Lys Arg Lys
180                               770                               775                               780
182 Asp Ala Met Gly Asn Thr Arg Gln Glu Met Thr His Met Val Asn Ala
183 785                               790                               795                               800
185 Met Asp Arg Ser Tyr Ala Asp Gln Ser Thr Leu His Ala Glu Asp Pro
186                               805                               810                               815
188 Leu Ser Leu Thr Phe Met Asp Gln His Asn Phe Ser Pro Arg Leu Pro
189                               820                               825                               830
191 Asn Asp Pro Leu Val Pro Thr Ala Val Leu Asp Glu Asn His Ser Ala
192                               835                               840                               845
194 Thr Ala Glu Ser Ser Arg Leu Leu Asp Val Pro Arg Tyr Leu Cys Glu
195                               850                               855                               860
197 Gly Thr Glu Ser Pro Tyr Gln Thr Gly Gln Leu His Pro Ala Ile Arg
198 865                               870                               875                               880
200 Val Ala Asp Leu Leu Gln His Ile Asn Leu Met Lys Thr Ser Asp Ser
201                               885                               890                               895
203 Tyr Gly Phe Lys Glu Glu Tyr Glu Ser Phe Phe Glu Gly Gln Ser Ala
204                               900                               905                               910
206 Ser Trp Asp Val Ala Lys Lys Asp Gln Asn Arg Ala Lys Asn Arg Tyr
207                               915                               920                               925
209 Gly Asn Ile Ile Ala Tyr Asp His Ser Arg Val Ile Leu Gln Pro Val
210                               930                               935                               940
212 Glu Asp Asp Pro Ser Ser Asp Tyr Ile Asn Ala Asn Tyr Ile Asp Ile

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213 945          950          955          960
215 Trp Leu Tyr Arg Asp Gly Tyr Gln Arg Pro Ser His Tyr Ile Ala Thr
216          965          970          975
218 Gln Gly Pro Val His Glu Thr Val Tyr Asp Phe Trp Arg Met Val Trp
219          980          985          990
221 Gln Glu Gln Ser Ala Cys Ile Val Met Val Thr Asn Leu Val Glu Val
222          995          1000          1005
224 Gly Arg Val Lys Cys Tyr Lys Tyr Trp Pro Asp Asp Thr Glu Val Tyr
225          1010          1015          1020
227 Gly Asp Phe Lys Val Thr Cys Val Glu Met Glu Pro Leu Ala Glu Tyr
228 1025          1030          1035          1040
230 Val Val Arg Thr Phe Thr Leu Glu Arg Arg Gly Tyr Asn Glu Ile Arg
231          1045          1050          1055
233 Glu Val Lys Gln Phe His Phe Thr Gly Trp Pro Asp His Gly Val Pro
234          1060          1065          1070
236 Tyr His Ala Thr Gly Leu Leu Ser Phe Ile Arg Arg Val Lys Leu Ser
237          1075          1080          1085
239 Asn Pro Pro Ser Ala Gly Pro Ile Val Val His Cys Ser Ala Gly Ala
240          1090          1095          1100
242 Gly Arg Thr Gly Cys Tyr Ile Val Ile Asp Ile Met Leu Asp Met Ala
243 1105          1110          1115          1120
245 Glu Arg Glu Gly Val Val Asp Ile Tyr Asn Cys Val Lys Ala Leu Arg
246          1125          1130          1135
248 Ser Arg Arg Ile Asn Met Val Gln Thr Glu Glu Gln Tyr Ile Phe Ile
249          1140          1145          1150
251 His Asp Ala Ile Leu Glu Ala Cys Leu Cys Gly Glu Thr Ala Ile Pro
252          1155          1160          1165
254 Val Cys Glu Phe Lys Ala Ala Tyr Phe Asp Met Ile Arg Ile Asp Ser
255          1170          1175          1180
257 Gln Thr Asn Ser Ser His Leu Lys Asp Glu Phe Gln Thr Leu Asn Ser
258 1185          1190          1195          1200
260 Val Thr Pro Arg Leu Gln Ala Glu Asp Cys Ser Ile Ala Cys Leu Pro
261          1205          1210          1215
263 Arg Asn His Asp Lys Asn Arg Phe Met Asp Met Leu Pro Pro Asp Arg
264          1220          1225          1230
266 Cys Leu Pro Phe Leu Ile Thr Ile Asp Gly Glu Ser Ser Asn Tyr Ile
267          1235          1240          1245
269 Asn Ala Ala Leu Met Asp Ser Tyr Arg Gln Pro Ala Ala Phe Ile Val
270          1250          1255          1260
272 Thr Gln Tyr Pro Leu Pro Asn Thr Val Lys Asp Phe Trp Arg Leu Val
273 1265          1270          1275          1280
275 Tyr Asp Tyr Gly Cys Thr Ser Ile Val Met Leu Asn Glu Val Asp Leu
276          1285          1290          1295
278 Ser Gln Gly Cys Pro Gln Tyr Trp Pro Glu Glu Gly Met Leu Arg Tyr
279          1300          1305          1310
281 Gly Pro Ile Gln Val Glu Cys Met Ser Cys Ser Met Asp Cys Asp Val
282          1315          1320          1325
284 Ile Asn Arg Ile Phe Arg Ile Cys Asn Leu Thr Arg Pro Gln Glu Gly
285          1330          1335          1340

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287 Tyr Leu Met Val Gln Gln Phe Gln Tyr Leu Gly Trp Ala Ser His Arg
288 1345          1350          1355          1360
290 Glu Val Pro Gly Ser Lys Arg Ser Phe Leu Lys Leu Ile Leu Gln Val
291          1365          1370          1375
293 Glu Lys Trp Gln Glu Glu Cys Glu Glu Gly Glu Gly Arg Thr Ile Ile
294          1380          1385          1390
296 His Cys Leu Asn Gly Gly Gly Arg Ser Gly Met Phe Cys Ala Ile Gly
297          1395          1400          1405
299 Ile Val Val Glu Met Val Lys Arg Gln Asn Val Val Asp Val Phe His
300          1410          1415          1420
302 Ala Val Lys Thr Leu Arg Asn Ser Lys Pro Asn Met Val Glu Ala Pro
303 1425          1430          1435          1440
305 Glu Gln Tyr Arg Phe Cys Tyr Asp Val Ala Leu Glu Tyr Leu Glu Ser
306          1445          1450          1455
308 Ser
311 <210> SEQ ID NO: 2
312 <211> LENGTH: 1439
313 <212> TYPE: PRT
314 <213> ORGANISM: Homo sapiens
316 <400> SEQUENCE: 2
317 Met Asp Thr Thr Ala Ala Ala Ala Leu Pro Ala Phe Val Ala Leu Leu
318 1          5          10          15
320 Leu Leu Ser Pro Trp Pro Leu Leu Gly Ser Ala Gln Gly Gln Phe Ser
321          20          25          30
323 Ala Gly Gly Cys Thr Phe Asp Asp Gly Pro Gly Ala Cys Asp Tyr His
324          35          40          45
326 Gln Asp Leu Tyr Asp Asp Phe Glu Trp Val His Val Ser Ala Gln Glu
327          50          55          60
329 Pro His Tyr Leu Pro Pro Glu Met Pro Gln Gly Ser Tyr Met Ile Val
330 65          70          75          80
332 Asp Ser Ser Asp His Asp Pro Gly Glu Lys Ala Arg Leu Gln Leu Pro
333          85          90          95
335 Thr Met Lys Glu Asn Asp Thr His Cys Ile Asp Phe Ser Tyr Leu Leu
336          100          105          110
338 Tyr Ser Gln Lys Gly Leu Asn Pro Gly Thr Leu Asn Ile Leu Val Arg
339          115          120          125
341 Val Asn Lys Gly Pro Leu Ala Asn Pro Ile Trp Asn Val Thr Gly Phe
342          130          135          140
344 Thr Gly Arg Asp Trp Leu Arg Ala Glu Leu Ala Val Ser Thr Phe Trp
345 145          150          155          160
347 Pro Asn Glu Tyr Gln Val Ile Phe Glu Ala Glu Val Ser Gly Gly Arg
348          165          170          175
350 Ser Gly Tyr Ile Ala Ile Asp Asp Ile Gln Val Leu Ser Tyr Pro Cys
351          180          185          190
353 Asp Lys Ser Pro His Phe Leu Arg Leu Gly Asp Val Glu Val Asn Ala
354          195          200          205
356 Gly Gln Asn Ala Thr Phe Gln Cys Ile Ala Thr Gly Arg Asp Ala Val
357          210          215          220
359 His Asn Lys Leu Trp Leu Gln Arg Arg Asn Gly Glu Asp Ile Pro Val

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**VERIFICATION SUMMARY**

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